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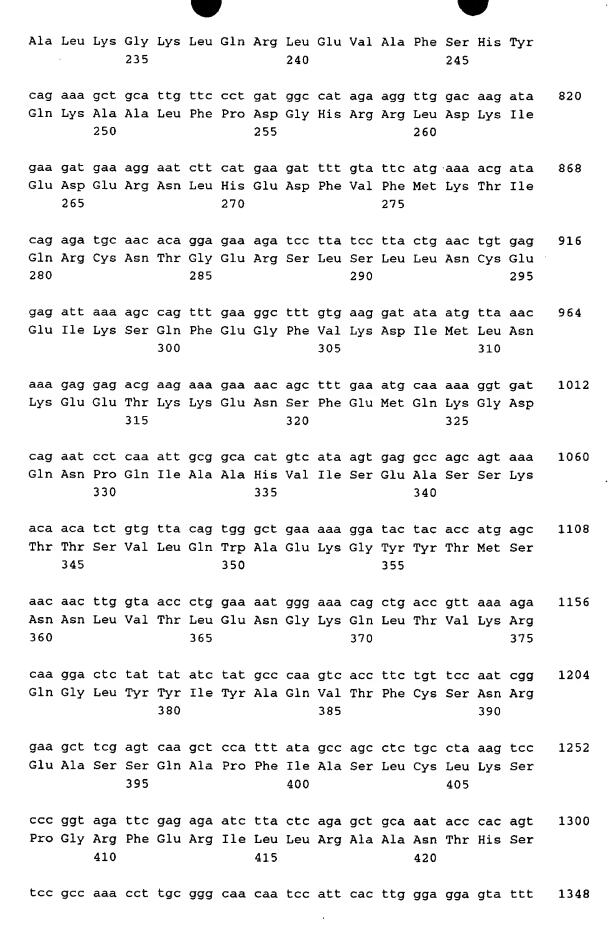
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125

120

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<301> Anderson, D M.
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<302> A homologue of the TNF receptor and its ligand enhance
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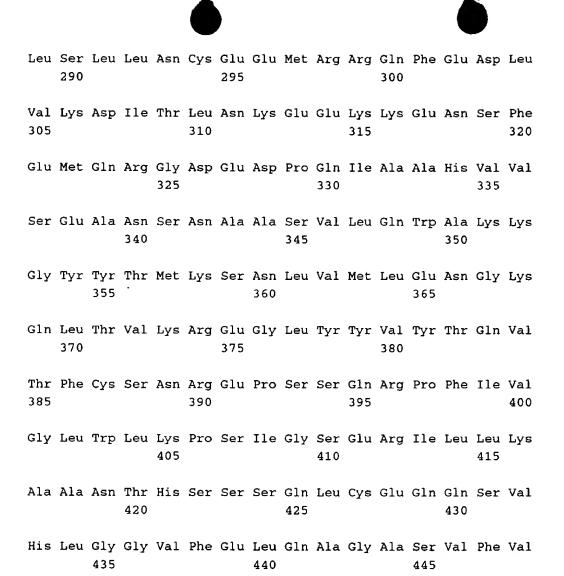
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<303> J. Exp. Med.
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cag aat cct caa att gcg gca cat gtc ata agt gag gcc agc agt aaa 1060 Gln Asn Pro Gln Ile Ala Ala His Val Ile Ser Glu Ala Ser Ser Lys 330 335 340

aca aca tct gtg tta cag tgg gct gaa aaa gga tac tac acc atg agc 1108 Thr Thr Ser Val Leu Gln Trp Ala Glu Lys Gly Tyr Tyr Thr Met Ser 345 350 355

aac aac ttg gta acc ctg gaa aat ggg aaa cag ctg acc gtt aaa aga 1156 Asn Asn Leu Val Thr Leu Glu Asn Gly Lys Gln Leu Thr Val Lys Arg 360 365 370 375

caa gga ctc tat tat atc tat gcc caa gtc acc ttc tgt tcc aat cgg 1204 Gln Gly Leu Tyr Tyr Ile Tyr Ala Gln Val Thr Phe Cys Ser Asn Arg 380 385 390

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ccc ggt aga ttc gag aga atc tta ctc aga gct gca aat acc cac agt 1300 Pro Gly Arg Phe Glu Arg Ile Leu Leu Arg Ala Ala Asn Thr His Ser 410 415 420

tcc gcc aaa cct tgc ggg caa caa tcc att cac ttg gga gga gta ttt 1348 Ser Ala Lys Pro Cys Gly Gln Gln Ser Ile His Leu Gly Gly Val Phe 425 430 435

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caa gtg age cat gge act gge tte acg tee ttt gge tta ete aaa ete 1444 Gln Val Ser His Gly Thr Gly Phe Thr Ser Phe Gly Leu Leu Lys Leu 460 465 470

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<213> Artificial Sequence

<223> Description of Artificial Sequence: Murine surfactant protein D (without the CRD) fused to the extracellular portion of human CD40L

260

275

265

280

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285

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Leu Ser Leu Leu Asn Cys Glu Glu Ile Lys Ser Gln Phe Glu Gly Phe
290 295 300

Val Lys Asp Ile Met Leu Asn Lys Glu Glu Thr Lys Lys Glu Asn Ser
305 310 315 320

Phe Glu Met Gln Lys Gly Asp Gln Asn Pro Gln Ile Ala Ala His Val 325 330 335

Ile Ser Glu Ala Ser Ser Lys Thr Thr Ser Val Leu Gln Trp Ala Glu 340 345 350

Lys Gly Tyr Tyr Thr Met Ser Asn Asn Leu Val Thr Leu Glu Asn Gly 355 360 365

Lys Gln Leu Thr Val Lys Arg Gln Gly Leu Tyr Tyr lle Tyr Ala Gln 370 375 380

Val Thr Phe Cys Ser Asn Arg Glu Ala Ser Ser Gln Ala Pro Phe Ile 385 390 395 400

Ala Ser Leu Cys Leu Lys Ser Pro Gly Arg Phe Glu Arg Ile Leu Leu 405 410 415

Arg Ala Asn Thr His Ser Ser Ala Lys Pro Cys Gly Gln Gln Ser 420 425 430

Ile His Leu Gly Gly Val Phe Glu Leu Gln Pro Gly Ala Ser Val Phe
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<300>
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   Guo, N
   Dowler, L L.
   Tauber, A I.
   Motwani, M
<302> Mouse surfactant protein-D. cDNA cloning,
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<303> J. Immunol.
<304> 155
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<301> Anderson, D M.
   Maraskovsky, E
   Billingsley, W L.
   Dougall, W C.
<302> A homologue of the TNF receptor and its ligand enhance
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<303> Nature
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<307> 1997
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ctt gtc ttg ctt gta cag ccc ctg gga aat ctg gga gca gaa atg aag 100

gcc tta aaa gga aaa cta cag cgt cta gag gtt gcc ttc tcc cac tat 772
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235 240 245

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aga ata tca gaa gac agc act cac tgc ttt tat aga atc ctg aga ctc 868
Arg Ile Ser Glu Asp Ser Thr His Cys Phe Tyr Arg Ile Leu Arg Leu
265 270 275

cat gaa aac gca ggt ttg cag gac tcg act ctg gag agt gaa gac aca 916 His Glu Asn Ala Gly Leu Gln Asp Ser Thr Leu Glu Ser Glu Asp Thr 280 285 290 295

cta cct gac tcc tgc agg agg atg aaa caa gcc ttt cag ggg gcc gtg 964 Leu Pro Asp Ser Cys Arg Arg Met Lys Gln Ala Phe Gln Gly Ala Val 300 305 310

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cca gct atg atg gaa ggc tca tgg ttg gat gtg gcc cag cga ggc aag 1060 Pro Ala Met Met Glu Gly Ser Trp Leu Asp Val Ala Gln Arg Gly Lys 330 335 340

cct gag gcc cag cca ttt gca cac ctc acc atc aat gct gcc agc atc 1108 Pro Glu Ala Gln Pro Phe Ala His Leu Thr Ile Asn Ala Ala Ser Ile 345 350 355

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cat cat gaa aca tcg gga agc gta cct aca gac tat ctt cag ctg atg 1300 His His Glu Thr Ser Gly Ser Val Pro Thr Asp Tyr Leu Gln Leu Met 410 415 420

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130

145

135

150

140

Glu Val Gly Ala Pro Gly Met Gln Gly Ser Thr Gly Ala Lys Gly Ser

160

155

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- Ile Pro Ser Ser His Asn Leu Met Lys Gly Gly Ser Thr Lys Asn Trp
  435 440 445

Thr Asp Tyr Leu Gln Leu Met Val Tyr Val Val Lys Thr Ser Ile Lys

Ser Gly Asn Ser Glu Phe His Phe Tyr Ser Ile Asn Val Gly Gly Phe 455 450 460 Phe Lys Leu Arg Ala Gly Glu Glu Ile Ser Ile Gln Val Ser Asn Pro 470 475 480 Ser Leu Leu Asp Pro Asp Gln Asp Ala Thr Tyr Phe Gly Ala Phe Lys 485 490 495 Val Gln Asp Ile Asp 500 <210>5 <211> 1477 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Murine surfactant protein D (except CRD) fused to the extracellular domain of murine CD40 ligand <220> <221> 5'UTR <222> (7)..(31) <223> 5' UTR from rat surfactant protein D <220> <221> sig\_peptide <222> (32)..(88) <223> Signal peptide from murine surfactant protein D <220> <221> CDS <222> (32)..(1441) <220> <221> misc\_recomb <222> (88)..(799) <223> Mature murine surfactant protein D including hub region, collagenous portion, and neck, but excluding carbohydrate recognition domain (CRD) <220> <221> misc\_feature <222>(800)..(1441) <223> Murine CD40 ligand extracellular region, including stalk <300> <301> Motwani, M

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   Guo, N
   Dowler, L L.
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   Motwani, M
<302> Mouse surfactant protein-D. cDNA cloning,
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<303> J. Immunol.
<304> 155
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<307> 1995
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   Sato, TA.
   Clifford, K N.
<302> Molecular and biological characterization of a murine
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Leu Val Leu Val Gln Pro Leu Gly Asn Leu Gly Ala Glu Met Lys
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Ser Leu Ser Gln Arg Ser Val Pro Asn Thr Cys Thr Leu Val Met Cys
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Ser Pro Thr Glu Asn Gly Leu Pro Gly Arg Asp Gly Arg Asp Gly Arg
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80

85

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1477

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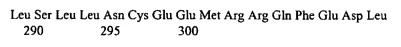
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<213> Artificial Sequence

<223> Description of Artificial Sequence: Murine surfactant protein D (except CRD) fused to the extracellular domain of murine CD40 ligand



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- Arg Asp Gly Arg Asp Gly Arg Glu Gly Pro Arg Gly Glu Lys Gly Asp 50 55 60
- Pro Gly Leu Pro Gly Pro Met Gly Leu Ser Gly Leu Gln Gly Pro Thr 65 70 75 80
- Gly Pro Val Gly Pro Lys Gly Glu Asn Gly Ser Ala Gly Glu Pro Gly 85 90 95
- Pro Lys Gly Glu Arg Gly Leu Ser Gly Pro Pro Gly Leu Pro Gly Ile 100 105 110
- Pro Gly Pro Ala Gly Lys Glu Gly Pro Ser Gly Lys Gln Gly Asn Ile 115 120 125
- Gly Pro Gln Gly Lys Pro Gly Pro Lys Gly Glu Ala Gly Pro Lys Gly 130 135 140
- Glu Val Gly Ala Pro Gly Met Gln Gly Ser Thr Gly Ala Lys Gly Ser 145 150 155 160
- Thr Gly Pro Lys Gly Glu Arg Gly Ala Pro Gly Val Gln Gly Ala Pro . 165 170 175
- Gly Asn Ala Gly Ala Ala Gly Pro Ala Gly Pro Ala Gly Pro Gln Gly 180 185 190
- Ala Pro Gly Ser Arg Gly Pro Pro Gly Leu Lys Gly Asp Arg Gly Val 195 200 205
- Pro Gly Asp Arg Gly Ile Lys Gly Glu Ser Gly Leu Pro Asp Ser Ala 210 215 220
- Ala Leu Arg Gln Gln Met Glu Ala Leu Lys Gly Lys Leu Gln Arg Leu 225 230 235 240
- Glu Val Ala Phe Ser His Tyr Gln Lys Ala Ala Leu Phe Pro Asp Gly 245 250 255
- His Arg Arg Leu Asp Lys Val Glu Glu Val Asn Leu His Glu Asp 260 265 270
- Phe Val Phe Ile Lys Lys Leu Lys Arg Cys Asn Lys Gly Glu Gly Ser 275 280 285



Val Lys Asp Ile Thr Leu Asn Lys Glu Glu Lys Lys Glu Asn Ser Phe 305 310 315 320

Glu Met Gln Arg Gly Asp Glu Asp Pro Gln Ile Ala Ala His Val Val 325 330 335

Ser Glu Ala Asn Ser Asn Ala Ala Ser Val Leu Gln Trp Ala Lys Lys 340 345 350

Gly Tyr Tyr Thr Met Lys Ser Asn Leu Val Met Leu Glu Asn Gly Lys 355 360 365

Gln Leu Thr Val Lys Arg Glu Gly Leu Tyr Tyr Val Tyr Thr Gln Val

Thr Phe Cys Ser Asn Arg Glu Pro Ser Ser Gln Arg Pro Phe Ile Val 385 390 395 400

Gly Leu Trp Leu Lys Pro Ser Ile Gly Ser Glu Arg Ile Leu Leu Lys 405 410 415

Ala Ala Asn Thr His Ser Ser Ser Gln Leu Cys Glu Gln Gln Ser Val 420 425 430

His Leu Gly Gly Val Phe Glu Leu Gln Ala Gly Ala Ser Val Phe Val 435 440 445

Asn Val Thr Glu Ala Ser Gln Val Ile His Arg Val Gly Phe Ser Ser 450 455 460

Phe Gly Leu Leu Lys Leu 465 470